

## Effect of salinity on chlorophyll concentration, leaf area, yield and yield components of rice genotypes grown under saline environment

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### Abstract

Salt tolerance in eighteen advanced rice genotypes was studied under an artificially salinized ( $EC=8.5 \text{ dSm}^{-1}$ ) soil conditions after 90 days of transplanting. The results showed that the yield per plant, chlorophyll concentrations, fertility percentage, and number of productive tillers, panicle length and number of primary braches per panicle of all the genotypes were reduced by salinity. However, genotypes viz. Jhona-349 x Basmati-370, NR-1, DM-59418, DM-63275, DM-64198 and DM-38-88 showed better salinity tolerance than others.

**Key words:** Rice, genotypes, salinity, dwarf mutant, chlorophyll

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### Introduction

Rice is the premier food grain crop of Pakistan for domestic consumption and export occupying an area of 2.2 million hectares (Zia, *et al.*, 1998). Soil salinity is one of the major constraints responsible for low agriculture production in Pakistan. Out of  $14.8 \times 10^6$  hectares irrigated land; 6 million hectares are salt affected. A loss of Rs. 20 million per year has been estimated on salt affected soils of Pakistan on account of decrease in agriculture production (Anonymous, 2003). Selection of salt tolerant cultivars is one of the most effective methods to increase the productivity of such soils. The major inhibitory effect of salinity on plant growth and yield has been attributed to: 1) osmotic effect 2) ion toxicity 3) nutritional imbalance leading to reduction in photosynthetic efficiency and other physiological disorders. Adverse effects of Salinity on seed germination and seedling growth as well as some physiological activities of cultivated plant species have been extensively investigated in Pakistan (Ashraf and Khan, 1993; Ashraf, *et al.*, 1991; Khan, *et al.*, 1995). Generally, the trend and magnitude of adverse changes varied with in species, varieties/genotypes according to the level of Salinization. So far little emphasis has been placed on aspects relevant to photosynthetic efficiency of plants at moderate and high salinities. It has been suggested that by increasing photosynthetic efficiency crop

production could be increased. Breeding for salinity tolerant in rice is difficult due to the involvement of several genes controlling the character and lack of sufficient knowledge of the mechanisms controlling salt tolerance (Aktita and Cabusley, 1988 and Yeo, *et al.*, 1990). Therefore the efforts have been made to develop the salt tolerant variety through induced mutations. Mutant lines used in this study are the derivatives of Basmati rice background having good quality grain with strong aroma. The aim of the present investigation is to provide information on the effect of salinity on chlorophyll concentration and yield and yield components of rice genotypes to see if there is any correlation between these variables.

### Materials and Methods

The study was conducted during the years 2001-02 in artificially salinized ( $6 \times 6 \times 1 \text{ m}$ ) concrete tanks located in net-house of Nuclear Institute for Agriculture and Biology (NIAB) Faisalabad (183 m above mean sea level 31 24 N and 730 05 E) Pakistan. Salinity was raised by mixing four commercial salts i.e.  $\text{Na}_2\text{SO}_4$ ,  $\text{NaCl}$ ,  $\text{MgCl}_2$  and  $\text{CaCl}_2$  in the ratio of 10:4:1:5 respectively on equivalent basis representing a type of salinity found in most parts of Pakistan (Qureshi, *et al.*, 1977). The experimental material comprised of eighteen varieties/variant/mutant lines (Table 1). The crop was also grown in normal soil

simultaneously. The soil was fertilized with urea at 185 kg/ka and DAP at 130 kg/ka. Four weeks old seedlings were transplanted with row to plant distance of 20 cm. in randomized complete block design. Ninety days after transplanting, three upper leaves of each tiller of three plants from each genotype were excised from each treatment and replication. Chlorophyll concentration (a, b and total) of these leaves, were determined according to Arnon (1949). At maturity the number of primary branches per panicle, number of productive tillers, panicle length and fertility percentage were also recorded in 6 genotypes, which showed tolerance against salinity. The harmful effects induced by salinity were computed in percent reduction over control (% ROC) with the following formula.

(% ROC) =

$$\frac{\text{Value in control} - \text{value in saline environment} \times 100}{\text{Value in control}}$$

Analysis of variance was applied to determine the significance of differences among the treatment and genotypes. Duncans Multiple New Range Test (DMRT) compared differences in mean at 5% probability (Steel and Torrie, 1980).

## Results

Yield per plant decreased significantly in response to salinity (Table 1) in all rice genotypes. The maximum yield under saline condition was recorded in DM-38-88 and NR-1 followed by others. When harmful effect of salinity was noted in the form of percent reduction over control, maximum reduction was observed in Super Basmati, KS-282 and Basmati-385 × NIAB-IRRI9 ranging from 48-50.6 %. DM-25 × NIAB-IRRI9, NIAB-IRRI-9 × DM-25, DM-59418, DM-38-88, DM-64198, Jhona-349 × Basmati-370, NIAB-Rice-1 and DM-63275 showed minimum reduction over control for yield per plant, which ranged from 15-36 %. The reduction in leaf area (Table 1) of all 18 rice genotypes under salinity stress plants has been attributed to suppressed cell division. Maximum percent reduction over control was noted in NIAB-IRRI-9, NIAB-Rice-1, DM-25 × NIAB-IRRI-9, DM-5-89 and super basmati while DM-63275, Jhona-349 × Basmati-370, DM-59418, DM-38-88, DM-64198 and Basmati-370 × NIAB-RICE-1 showed minimum percent reduction for leaf area. The biosynthesis of pigment fractions (chlorophyll a, b and total) was affected with salinity stress (Table 2).

Table 1: Effect of salinity on grain yield per plant and flag leaf area of different rice genotypes

Genotypes	Yield per plant (gms)			Flag leaf area (cm. <sup>2</sup> )		
	Control	EC=8.5d Sm <sup>-1</sup>	Percentage Reduction over Control	Control	EC=8.5d Sm <sup>-1</sup>	Percentage reduction over control
DM-63275	21.80667	13.91667	36.18159 CDE	43.88333	40.1	8.621338 D
NIAB-IRRI-9	19.56333	13.48333	31.07855 DEF	38.62333	20.33333	47.3548 A
KS-282	16.70667	8.323333	50.17958 A	40.13333	31.28333	22.0515 C
NIAB-RICE-1	20.27667	14.41667	28.90021 EF	80.89333	46.1	43.01137 A
DM-25xNIAB-IRRI-9	12.89333	10.91	15.3826 G	81.73333	40	51.06036 A
Jhona-349	20.55	11.44667	44.29844 AB	50.65	37.91	25.15301 BC
DM-5-89	16.90667	10.52667	37.73659 BCD	61.23333	31.37667	48.75884 A
NIAB-IRRI-9xDM-25	13.28333	9.666667	27.22708 F	72.73667	52.56667	27.73017 BC
Jhona-349xBasmati-370	18.46667	12.8	30.68593 DEF	48.3	39.67	17.86749 CD
Basmati-370xJhona-349	17.56667	11.94333	32.01142 DEF	48.43333	34.59333	28.57536 BC
Basmati-370	14.16333	7.873333	44.41044 AB	43.19333	31.32667	27.47336 BC
DM-59418	20.53333	14.73	28.26298 F	47.86333	39.51667	17.43853 CD
Basmati-385xNIAB-IRRI-9	16.96	8.75	48.40802 A	48.86333	32.78333	32.90811 B
Super Basmati	17.83333	8.81	50.59812 A	62.1	32.63333	47.45035 A
DM-38-88	20.30667	14.73333	27.44586 F	52	41.10667	20.94871 C
DM-64198	19.51333	14.25	26.973 F	52.51667	42.86667	18.37512 CD
Basmati-370xNIAB-RICE-1	18.09333	11.88333	34.32204 CDEF	50.42667	40.36667	19.94976 C
DM-3-89	17.36667	10.36333	40.32633 BC	59.41333	44.68	24.79802 BC

Mean in the same column sharing the letters did not differ significantly according to DMRT (P=0.05)

Table 2: Chlorophyll concentrations (a, b and total) of different rice genotypes grown under normal and saline environments

Genotypes	Chlorophyll (a)			Chlorophyll (b)			Chlorophyll (Total)		
	Control	EC=8.5d Sm <sup>-1</sup>	Percentage reduction over control	Control	EC=8.5d Sm <sup>-1</sup>	Percentage reduction over control	Control	EC=8.5d Sm <sup>-1</sup>	Percentage reduction over control
DM-63275	0.540	0.389	27.97 C	0.283	0.257	9.187279 EFG	0.822567	0.645667	21.507 CDEF
NIAB-IRRI-9	0.596	0.440	26.22 CD	0.365	0.290	20.54795 AB	0.938767	0.721267	22.84145 BCDE
KS-282	0.493	0.358	27.34 C	0.288	0.260	9.722222 EFG	0.7809	0.618033	20.85663 DEF
NIAB-RICE-1	0.479	0.441	7.92 G	0.280	0.256	8.571429 FGH	0.758867	0.697933	8.012744 I
DM-25xNIAB-IRRI-9	0.559	0.412	26.44 CD	0.299	0.280	6.354515 GH	0.8586	0.6912	19.49985 FG
Jhona-349	0.523	0.392	24.99 B	0.291	0.278	4.467354 H	0.811133	0.6704	17.35333 GH
DM-5-89	0.501	0.349	30.38 B	0.301	0.252	16.27907 BCD	0.8019	0.601333	25.01539 AB
NIAB-IRRI-9xDM-25	0.531	0.420	20.90 E	0.279	0.251	10.03584 EFG	0.810067	0.671333	17.12694 GH
Jhona-349xBasmati-370	0.544	0.449	17.52 F	0.320	0.278	13.125 DE	0.864	0.7273	15.8243 H
Basmati-370xJhona-349	0.572	0.411	28.22 C	0.317	0.261	17.66562 BC	0.889533	0.6717	24.49202 BC
Basmati-370	0.528	0.366	30.67 B	0.289	0.238	17.64706 BC	0.8172	0.604867	25.9863 AB
DM-59418	0.556	0.405	27.20 C	0.309	0.262	15.21036 CD	0.866167	0.6672	22.97811 BCDE
Basmati-385xNIAB-IRRI-9	0.491	0.370	24.70 D	0.300	0.231	23 A	0.791433	0.601267	24.03662 BCD
Super Basmati	0.516	0.376	27.24 C	0.273	0.257	5.860806 GH	0.789333	0.632567	19.85444 EFG
DM-38-88	0.560	0.392	30.01 B	0.292	0.255	12.67123 DEF	0.8513	0.6468	24.01885 BCD
DM-64198	0.602	0.404	32.89 A	0.320	0.261	18.4375 BC	0.921933	0.665033	27.86813 A
Basmati-370xNIAB-RICE-1	0.550	0.379	31.17 AB	0.301	0.277	7.973422 GH	0.5644	0.434067	15.39454 BCDE
DM-3-89	0.477	0.449	5.87 H	0.277	0.258	6.859206 GH	0.498833	0.467667	4.165616 I

Mean in the same column sharing the letters did not differ significantly according to DMRT (P<0.05)

Table 3: Influence of salinity on (a) number of primary branches per panicle (b) number of productive tillers (c) panicle length (d) fertility percentage of rice genotypes

Name of Genotypes	Salinity level d Sm <sup>-1</sup>		Percentage increase over control
	Control (EC = 1.80)	EC = 8.5	
DM-59418	a-10.5	8.6	18.09
	b-10.38	08.03	22.64
	c-35.52	28.97	18.44
	d-91.88	70.42	23.36
DM-64198	a-11.1	8.7	21.62
	b-11.86	08.35	29.68
	c-24.76	22.67	08.44
	d-97.00	84.59	12.79
Jhona-349 x Basmati-370	a-12.0	9.4	21.66
	b-11.24	08.40	25.27
	c-28.56	22.77	20.27
	d-94.04	85.53	09.05
DM-63275	a-10.95	8.8	19.63
	b-13.90	08.00	42.45
	c-31.34	27.80	11.01
	d-91.90	69.35	24.54
NIAB-RICE-1	a-12.2	9.8	19.67
	b-08.87	06.27	29.31
	c-29.73	25.67	13.66
	d-87.73	74.19	15.43
DM-38-88	a-12.3	10.2	17.07
	b-08.48	05.92	30.19
	c-29.95	25.40	15.19
	d-84.30	71.01	15.77

DM = Dwarf mutant

DM-64198, Basmati-370 and DM-5-89 showed maximum percent reduction over control for chlorophyll total concentration and were graded as sensitive to salinity stress. Genotypes DM-3-89, NIAB-Rice-1, Jhona-349 × Basmati-370 and NIAB-IRRI9 × DM-25 minimum reduction in chlorophyll concentrations and were graded as salt tolerant. Influence of salinity on yield components i.e. number of primary braches per panicle, number of productive tillers per plant, panicle length and fertility percentage is given in (Table 3). Results indicated that percent increase over control for number of primary braches per panicle ranged from 17.07-21.66, for number of productive tillers per plants 22.64–42.45, panicle length (cm.) 8.44-20.27 and fertility percentage ranged from 9.05-24.54 in six genotypes tested. All the six genotypes were graded as salt tolerant with respect to yield components.

### Discussion and Conclusion

When plants are grown under saline conditions, as soon as the new cell starts its

elongation process, the excess of salts modifies the metabolic activities of the cell wall causing the deposition of various materials which limit the cell wall elasticity. Secondary cell wall sooner, cell walls become rigid and consequently the turgor pressure efficiency in cell enlargement is decreased. The other expected causes of the reduction in yield per plant, leaf area and yield components in rice could be the shrinkage of the cell contents, reduced development and differentiation of tissues, unbalanced nutrition, damage of membrane and disturbed avoidance mechanism. The reduction in leaf area, yield and yield components under saline conditions were also due to reduced growth as a result of decreased water uptake, toxicity of sodium and chloride in the shoot cell as well as reduced photosynthesis. Reduction in chlorophyll concentrations is probably due to the inhibitory effect of the accumulated ions of various salts on the biosynthesis of the different chlorophyll fractions. Salinity affects the strength of the forces bringing the complex pigment protein-

liquid, in the chloroplast structure. As the chloroplast in membrane bound its stability is dependent on the membrane stability which under high salinity condition seldom remains intact due to which reduction in chlorophyll was recorded. Salt tolerance is not a function of single organ or plant attribute, but it is the product of all the plant attributes. Therefore a genotype exhibiting relative salt tolerance for all the plant attributes may be ideal one. Fortunately the mutants studied viz. DM-59418, DM-64198, and Jhona-349 × Basmati-370, DM-63275, NIAB-Rice-1 and DM-38-88 has shown comparatively minimum salinity induced reduction for the plant attributes. In this study some genotypes showed tolerance to salinity for the plant attributes that 6 genotypes lacked. The genotypes could be used as donors for these further improvements of mutant lines to establish definite relation with yield, chlorophyll concentration and leaf area. Further study would be initiated with this basic information. By using these mutant lines in breeding Programmed an improved ideotype of rice having higher chlorophyll concentration, more leaf area, early and better yield potential will be selected. This genotype possessing salt tolerance character will help in boosting up rice production in salt-affected soils. Therefore a genotype exhibiting relative salt tolerance for all the plant attributes may be ideal one. Fortunately the mutants studied viz. DM-59418, DM-64198, and Jhona-349 × Basmati-370, DM-63275, NIAB-Rice-1 and DM-38-88 has shown comparatively minimum salinity induced reduction for the plant attributes. In this study some genotypes showed tolerance to salinity for the plant attributes that 6 genotypes lacked. The genotypes could be used as donors for these further improvements of mutant lines to establish definite relation with yield, chlorophyll concentration and leaf area. Further study would be initiated with this basic information. By using these mutant lines in breeding Programmed an improved ideotype of rice having higher chlorophyll concentration, more leaf area, early and better yield potential will be selected. This genotype possessing salt tolerance character will help in boosting up rice production in salt-affected soils.

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